

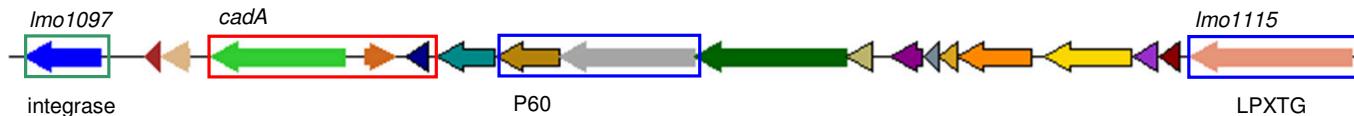
## Content and conservation of transposons ICELm1 and TN554 in species *L. monocytogenes*

Based on a homology cutoff >60% amino acid identity and >80% coverage. A black border denotes a deviation from the average codon usage of the chromosome.



### A) ICELm1:

1. *Listeria monocytogenes* 1-2a EGD-e refseq  
1131k .. 1152k (20917 bp)



### B) TN554:

1. *Listeria monocytogenes* 3c SLCC2479  
2765k .. 2776k (10476 bp)

2. *Listeria monocytogenes* 1-2c SLCC2372  
2766k .. 2776k (10476 bp)

